1641

RAW SEQUENCE LISTING DATE: 02/21/2001 PATENT APPLICATION: US/09/653,755A TIME: 13:16:23

Input Set : A:\Mab4gl0.app

Output Set: N:\CRF3\02212001\I653755A.raw



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3 <110> APPLICANT: Eisinger, Dominic P.
      4
              Stiles, Lynn
      5
              LaMarche, Arthur
              Jelinek, Thomas
      8 <120> TITLE OF INVENTION: Recombinant Monoclonal Antibody Specific for
              Phosphotyrosine-Containing Proteins
     11 <130> FILE REFERENCE: 724650-3
C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/653,755A
     14 <141> CURRENT FILING DATE: 2000-09-01
     16 <160> NUMBER OF SEQ ID NOS: 12
     18 <170> SOFTWARE: PatentIn Ver. 2.1
     20 <210> SEQ ID NO: 1
     21 <211> LENGTH: 1365
     22 <212> TYPE: DNA
     23 <213> ORGANISM: Artificial Sequence
     25 <220> FEATURE:
     26 <223> OTHER INFORMATION: Description of Artificial Sequence:cDNA for heavy
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     31 teetgeagga ettetgeata cacatteaet gaaaacaeeg tgeaetgggt gaageagage 120
     32 catggagaga gccttgagtg gattggaggt attaatcctt actatggtgg ttctatcttc 180
     33 agcccgaagt tcaagggcaa ggccacattg actgtagaca agtcctccag cacagcctac 240
     34 atggagetee geageetgae atetgaggat tetgeagtet attactgtge aagaaggget 300
     35 ggggcgtact actttgacta ctggggccaa ggcaccactc tcacagtctc ctcagccaaa 360
     36 acaacaccc catcagtcta tocactggcc cctgggtgtg gagatacaac tggttcctcc 420
     37 gtgactetgg gatgeetggt caagggetae tteeetgagt cagtgaetgt gaettggaae 480
     38 totggatoco tgtocagoag tgtgoacaco ttoccagoto teotgoagto tggactotác 540
     39 actatgagea geteagtgae tgteeeetee ageaeetgge eaagteagae egteaeetge 600
     40 agegttgete acceageeag cageaceaeg gtggacaaaa aacttgagee cagegggeee 660
    41 atttcaacaa tcaacccctg tcctccatgc aaggagtgtc acaaatgccc agctcctaac 720
    42 ctcgagggtg gaccatccgt cttcatcttc cctccaaata tcaaggatgt actcatgatc 780
    43 teectgacae ecaaggteae gtgtgtggtg gtggatgtga gegaggatga eceagaegte 840
    44 cagatcagct ggtttgtgaa caacgtggaa gtacacacag ctcagacaca aacccataga 900
    45 gaggattaca acagtactat ccgggtggtc agcaccetcc ccatccagca ccaggactgg 960
    46 atgagtggca aggagttcaa atgcaaggtc aacaacaaag acctcccatc acccatcgag 1020
    47 agaaccatct caaaaattaa agggctagtc agagctccac aagtatacat cttgccgcca 1080
    48 ccagcagage agttgtccag gaaagatgte agteteaett geetggtegt gggetteaac 1140
    49 cctggagaca tcagtgtgga gtggaccagc aatgggcata cagaggagaa ctacaaggac 1200
    50 accgcaccag teetggacte tgacggttet tactteatat atagcaaget caatatgaaa 1260
    51 acaagcaagt gggagaaaac agatteette teatgeaacg tgagacaega gggtetgaaa 1320
    52 aattactacc tgaagaagac catctcccgg tctccgggta aatga
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    56 <211> LENGTH: 645
    57 <212> TYPE: DNA
    58 <213> ORGANISM: Artificial Sequence
    60 <220> FEATURE:
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61 <223> OTHER INFORMATION: Description of Artificial Sequence:cDNA for light
         chain of recombinant antibody
64 <400> SEQUENCE: 2
65 gaaaatgtgc tcacccagtc tccagcaatc atgtctgcat ctccagggga aaaggtcacc 60
66 atgacctgca gggccagctc aagtgtaagt tecagttact tgcactggta teggcagaag 120
67 traggtgect coccaaact otggatttat agcacatoca acttggctto tggagtcoot 180
68 gctcgcttca gtggcagtgg gtctgggacc tcttactctc tcacaatcag cagtgtggag 240
69 gctgaagatg ctgccactta ttactgccag cagtacagtg gttaccggac gttcggtgga 300
70 ggcaccaagc tggaaatcaa acgggctgat gctgcaccaa ctgtatccat cttcccacca 360
71 tocagtgage agttaacate tggaggtgee teagtegtgt gettettgaa caacttetae 420
72 occagagaca tcaatgtcaa gtggaagatt gatggcagtg aacgacaaaa tggtgtcctg 480
73 aacagttgga ctgatcagga cagcaaagac agcacctaca gcatgagcag caccctcaca 540
74 ttgaccaagg acgagtatga acgacataac agctatacct gtgaggccac tcacaagaca 600
75 tcaacttcac ccatcgtcaa gagcttcaac aggaatgagt gttag .
78 <210> SEQ ID NO: 3
79 <211> LENGTH: 1389
80 <212> TYPE: DNA
81 <213> ORGANISM: Artificial Sequence
83 <220> FEATURE:
84 <223> OTHER INFORMATION: Description of Artificial Sequence:cDNA for heavy
         chain of recombinant antibody with 3'-histidine
86
         tag sequence
88 <400> SEQUENCE: 3
89 gaggtccagc tgcarcagtc tggacctgaa ctggtgaagc ctggggcttc agtgatgata 60
90 teetgeagga ettetgeata cacatteact gaaaacaeeg tgeactgggt gaageagage 120
91 catggagaga gccttgagtg gattggaggt attaatcctt actatggtgg ttctatcttc 180
92 agcccgaagt tcaagggcaa ggccacattg actgtagaca agtcctccag cacagcctac 240
93 atggagetee geageetgae atetgaggat tetgeagtet attactgtge aagaaggget 300
94 ggggcgtact actttgacta ctggggccaa ggcaccactc tcacagtctc ctcagccaaa 360
95 acaacacece cateagteta tecaetggee eetgggtgtg gagatacaae tggtteetee 420
96 gtgactctgg gatgcctggt caagggctac ttccctgagt cagtgactgt gacttggaac 480
97 tetggatece tgtecageag tgtgeacace tteecagete teetgeagte tggaetetae 540
98 actatgagca geteagtgac tgtecectec ageaeetgge caagteagae egteaeetge 600
99 agogttgctc acccagecag cagcaccacg gtggacaaaa aacttgagec cagegggeec 660
100 attteaacaa teaacceetg teeteeatge aaggagtgte acaaatgeee ageteetaac 720
101 ctcgagggtg gaccatccgt cttcatcttc cctccaaata tcaaggatgt actcatgatc 780
102 tecetgacae ecaaggteae gtgtgtggtg gtggatgtga gegaggatga eccagaegte 840
103 cagatcaget ggtttgtgaa caacgtggaa gtacacacag etcagacaca aacccataga 900
104 gaggattaca acagtactat cogggtggtc agcaccotoc coatocagca coaggactgg 960
105 atgagtggca aggagttcaa atgcaaggtc aacaacaaag acctcccatc acccatcgag 1020
106 agaaccatct caaaaattaa agggctagtc agagctccac aagtatacat cttgccgcca 1080
107 ccagcagage agttgtccag gaaagatgte agteteaett geetggtegt gggetteaae 1140
108 cctggagaca tcagtgtgga gtggaccagc aatgggcata cagaggagaa ctacaaggac 1200
109 accgcaccag teetggacte tgaeggttet taetteatat atagcaaget caatatgaaa 1260
110 acaagcaagt gggagaaaac agatteette teatgcaacg tgagacacga gggtetgaaa 1320
111 aattactacc tgaagaagac catctcccgg tctccgggta aaggtggcca tcaccaccat 1380
112 caccattga
115 <210> SEQ ID NO: 4
116 <211> LENGTH: 454
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117 <212> TYPE: PRT
118 <213> ORGANISM: Artificial Sequence
120 <220> FEATURE:
121 <223> OTHER INFORMATION: Description of Artificial Sequence: Amino acid
         sequence for heavy chain of recombinant antibody
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128 Ser Val Met Ile Ser Cys Arg Thr Ser Ala Tyr Thr Phe Thr Glu Asn
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                                    2.5
                                                       30
131 Thr Val His Trp Val Lys Gln Ser His Gly Glu Ser Leu Glu Trp Ile
132
           35
                                40
134 Gly Gly Ile Asn Pro Tyr Tyr Gly Gly Ser Ile Phe Ser Pro Lys Phe
                            55
137 Lys Gly Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser Thr Ala Tyr
138 65
                        70
140 Met Glu Leu Arg Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys
                    8.5
                                       90
143 Ala Arg Arg Ala Gly Ala Tyr Tyr Phe Asp Tyr Trp Gly Gln Gly Thr
              100
                                  105
146 Thr Leu Thr Val Ser Ser Ala Lys Thr Thr Pro Pro Ser Val Tyr Pro
147 115
                               120
                                          125
149 Leu Ala Pro Gly Cys Gly Asp Thr Thr Gly Ser Ser Val Thr Leu Gly
                          135
                                       140
152 Cys Leu Val Lys Gly Tyr Phe Pro Glu Ser Val Thr Val Thr Trp Asn
                       150
                                           155
153 145
155 Ser Gly Ser Leu Ser Ser Ser Val His Thr Phe Pro Ala Leu Leu Gln
                                       170
158 Ser Gly Leu Tyr Thr Met Ser Ser Val Thr Val Pro Ser Ser Thr
               180
                                   185
                                                      190
161 Trp Pro Ser Gln Thr Val Thr Cys Ser Val Ala His Pro Ala Ser Ser
         195
                               200
164 Thr Thr Val Asp Lys Leu Glu Pro Ser Gly Pro Ile Ser Thr Ile
                           215
167 Asn Pro Cys Pro Pro Cys Lys Glu Cys His Lys Cys Pro Ala Pro Asn
                       230
                                           235
170 Leu Glu Gly Gly Pro Ser Val Phe Ile Phe Pro Pro Asn Ile Lys Asp
                               250
                   245
                                                          255
173 Val Leu Met Ile Ser Leu Thr Pro Lys Val Thr Cys Val Val Asp
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                                                       270
                                   265
176 Val Ser Glu Asp Asp Pro Asp Val Gln Ile Ser Trp Phe Val Asn Asn
                               280
           275
179 Val Glu Val His Thr Ala Gln Thr Gln Thr His Arg Glu Asp Tyr Asn
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                           295
                                               300
182 Ser Thr Ile Arg Val Val Ser Thr Leu Pro Ile Gln His Gln Asp Trp
                      310
                                          315
185 Met Ser Gly Lys Glu Phe Lys Cys Lys Val Asn Asn Lys Asp Leu Pro
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                                      330
188 Ser Pro Ile Glu Arg Thr Ile Ser Lys Ile Lys Gly Leu Val Arg Ala
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189
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191 Pro Gln Val Tyr Ile Leu Pro Pro Pro Ala Glu Gln Leu Ser Arg Lys
192 355
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194 Asp Val Ser Leu Thr Cys Leu Val Val Gly Phe Asn Pro Gly Asp Ile
                           375
        370
197 Ser Val Glu Trp Thr Ser Asn Gly His Thr Glu Glu Asn Tyr Lys Asp
                       390
                                           395
200 Thr Ala Pro Val Leu Asp Ser Asp Gly Ser Tyr Phe Ile Tyr Ser Lys
                    405
                                       410
203 Leu Asn Met Lys Thr Ser Lys Trp Glu Lys Thr Asp Ser Phe Ser Cys
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206 Asn Val Arg His Glu Gly Leu Lys Asn Tyr Tyr Leu Lys Lys Thr Ile
207 435
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209 Ser Arg Ser Pro Gly Lys
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213 <210> SEQ ID NO: 5
214 <211> LENGTH: 214
215 <212> TYPE: PRT
216 <213> ORGANISM: Artificial Sequence
218 <220> FEATURE:
219 <223> OTHER INFORMATION: Description of Artificial Sequence: Amino acid
        sequence for light chain of recombinant antibody
222 <400> SEQUENCE: 5
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226 Glu Lys Val Thr Met Thr Cys Arg Ala Ser Ser Ser Val Ser Ser Ser
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                                                       30
229 Tyr Leu His Trp Tyr Arg Gln Lys Ser Gly Ala Ser Pro Lys Leu Trp
                                4.0
232 Ile Tyr Ser Thr Ser Asn Leu Ala Ser Gly Val Pro Ala Arg Phe Ser
233
       50
                            55
                                                60
235 Gly Ser Gly Ser Gly Thr Ser Tyr Ser Leu Thr Ile Ser Ser Val Glu
                        7.0
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238 Ala Glu Asp. Ala Ala Thr Tyr Tyr Cys Gln Gln Tyr Ser Gly Tyr Arg
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241 Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Arg Ala Asp Ala Ala
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244 Pro Thr Val Ser Ile Phe Pro Pro Ser Ser Glu Gln Leu Thr Ser Gly
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247 Gly Ala Ser Val Val Cys Phe Leu Asn Asn Phe Tyr Pro Arg Asp Ile
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250~\mathrm{Asn} Val Lys Trp Lys Ile Asp Gly Ser Glu Arg Gln Asn Gly Val Leu
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253 Asn Ser Trp Thr Asp Gln Asp Ser Lys Asp Ser Thr Tyr Ser Met Ser
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                                       170
256 Ser Thr Leu Thr Leu Thr Lys Asp Glu Tyr Glu Arg His Asn Ser Tyr
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                                  185
259 Thr Cys Glu Ala Thr His Lys Thr Ser Thr Ser Pro Ile Val Lys Ser
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266 <210> SEQ ID NO: 6
267 <211> LENGTH: 462
268 <212> TYPE: PRT
269 <213> ORGANISM: Artificial Sequence
271 <220> FEATURE:
272 <223> OTHER INFORMATION: Description of Artificial Sequence: Amino acid
         sequence for heavy chain of recombinant antibody
         with C-terminal histidine tag sequence
274
276 <400> SEQUENCE: 6
277 Glu Val Gln Leu Gln Gln Ser Gly Pro Glu Leu Val Lys Pro Gly Ala
    1
280 Ser Val Met Ile Ser Cys Arg Thr Ser Ala Tyr Thr Phe Thr Glu Asn
281
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                                    25
                                                        3.0
283 Thr Val His Trp Val Lys Gln Ser His Gly Glu Ser Leu Glu Trp Ile
284 35
                                40
286 Gly Gly Ile Asn Pro Tyr Tyr Gly Gly Ser Ile Phe Ser Pro Lys Phe
287 50
                            55
                                                60
289 Lys Gly Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser Thr Ala Tyr
                        70
                                            75
292 Met Glu Leu Arg Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys
                    85
                                        90
                                                            95
295 Ala Arg Arg Ala Gly Ala Tyr Tyr Phe Asp Tyr Trp Gly Gln Gly Thr
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296
               100
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298 Thr Leu Thr Val Ser Ser Ala Lys Thr Thr Pro Pro Ser Val Tyr Pro
299
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                               120
301 Leu Ala Pro Gly Cys Gly Asp Thr Thr Gly Ser Ser Val Thr Leu Gly
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304 Cys Leu Val Lys Gly Tyr Phe Pro Glu Ser Val Thr Val Thr Trp Asn
305 145
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307 Ser Gly Ser Leu Ser Ser Ser Val His Thr Phe Pro Ala Leu Leu Gln
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                   165
310 Ser Gly Leu Tyr Thr Met Ser Ser Val Thr 'Val Pro Ser Ser Thr
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                                   185
313 Trp Pro Ser Gln Thr Val Thr Cys Ser Val Ala His Pro Ala Ser Ser
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                                                   205
316 Thr Thr Val Asp Lys Lys Leu Glu Pro Ser Gly Pro Ile Ser Thr Ile
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                                               220
319 Asn Pro Cys Pro Pro Cys Lys Glu Cys His Lys Cys Pro Ala Pro Asn
                       230
                                           235
322 Leu Glu Gly Gly Pro Ser Val Phe Ile Phe Pro Pro Asn Ile Lys Asp
                                       250
                   245
325 Val Leu Met Ile Ser Leu Thr Pro Lys Val Thr Cys Val Val Val Asp
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                                   265
                                                       270
328 Val Ser Glu Asp Asp Pro Asp Val Gln Ile Ser Trp Phe Val Asn Asn
                               280
    275
331 Val Glu Val His Thr Ala Gln Thr Gln Thr His Arg Glu Asp Tyr Asn
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VERIFICATION SUMMARYDATE: 02/21/2001PATENT APPLICATION:US/09/653,755ATIME: 13:16:24

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L:13 M:270 C: Current Application Number differs, Replaced Current Application Number